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pH-controlled liposomes for enhanced cell penetration in tumor environment

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ABSTRACT

An innovative pH-switchable colloidal system that can be exploited for site-selective anticancer drug delivery has been generated by liposomes decoration with a new novel synthetic non-peptidic oligo-Arginine cell-penetration enhancer (CPE) and a quenching PEGylated counterpart that detaches from the vesicle surface under the acidic conditions of tumors. The CPE module (*Arg₄-DAG*) is formed by four arginine units conjugated to a first generation (G1) 2,2-bis(hydroxymethyl)propionic acid (bis-MPA)/2,2-bis(aminomethyl)propionic acid (bis-AMPA) polyester dendron terminating with 1,2-distearoyl-3-azidopropane for liposome bilayer insertion. The zeta potential of the *Arg₄-DAG* -decorated liposomes increased up to +32 mV as the *Arg₄-DAG*/lipids molar ratio increased. The *Arg₄-DAG* liposome shielding at pH 7.4 was provided by methoxy-PEG_{5kDa}-polymethacryloyl sulfadimethoxine (mPEG_{5kDa}-SDM₈) with 7.1 apparent pKa. Zeta potential, surface plasmon resonance and synchrotron small-angle X-ray scattering analyses showed that at pH 7.4 mPEG_{5kDa}-SDM₈ associates with polycationic *Arg₄-DAG*-decorated liposomes yielding liposomes with neutral zeta potential. At pH 6.5, which mimics the tumor environment, mPEG_{5kDa}-SDM₈ detaches from the liposome surface yielding *Arg₄-DAG* exposure. Flow cytometry and confocal microscopy showed a 30-fold higher HeLa cancer cell association of the *Arg₄-DAG*-decorated liposomes compared to non-decorated liposomes. At pH 7.4, the mPEG_{5kDa}-SDM₈ coated liposomes undergo low cell association while remarkable cell association occurred at pH 6.5, which allowed for the controlled intracellular delivery of model macromolecules and small molecules loaded in the liposome under tumor conditions.

1. INTRODUCTION

New generation high-performance drug delivery colloidal systems have been developed to increase the tumor targeting and reduce off-target toxicity of chemotherapeutic agents.¹⁻² These systems are often designed to exploit the peculiar physio-pathological features at the tumor site to timely and spatially control the drug release.³⁻⁴

Targeting agents have been widely used to decorate drug nanocarriers for selective biorecognition of cells and intracellular delivery of therapeutic agents. However, biorecognition approaches may fail because tumor cell populations are extremely heterogeneous and expression levels of membrane targets can significantly vary.⁵⁻⁶ Furthermore, multivalent targeted nanocarriers may affect cellular receptor recycling efficiency during endocytosis, which limits further cargo internalization.⁷

A different approach for intracellular delivery of colloidal vehicles of anticancer drugs exploits cell penetration peptides (CPPs). Cell penetrating peptides, namely natural and synthetic TAT (transactivator of transcription)-derived peptides,⁸ have been exploited for intracellular delivery of a variety of colloidal cargoes, including proteins, oligonucleotides, nanovesicles and nanoparticles.⁹⁻¹⁰ Although a general consensus on the uptake mechanism of CPPs has not yet been achieved, two major cellular mechanisms of endocytic and non-endocytic pathways have been proposed. These mechanisms have been reported to depend on the transducing peptide sequences, the biophysical characteristics of the cargo and the cell type-dependent composition of the plasma membrane.¹¹⁻¹² However, high positive charge density resulting from protonation of the basic amino acid residues including arginine (Arg), lysine (Lys) and histidine (His) has been shown to be a structural requisite for CPP activity. Indeed, high positively charged amino

acid sequences are engaged providing charge-to-charge interactions with anionic proteoglycans, in particular heparin sulfate proteoglycans (HSPGs), on the cell membrane corona.¹² Brock *et al* reported that the cell up-take efficiency of CPPs increases with the decrease of the CPPs/HSPGs molar ratio and with the increase of the local Arg density in the peptide sequence.¹³⁻¹⁴

The guanidinium groups of Arg in the CPP sequences are structurally planar “Y” shaped molecules with delocalized positive charges, which allows for both electrostatic and hydrogen bonding with anionic and polar molecules.¹⁵ Accordingly, Arg interact through bi- or multi-dentate hydrogen bonds with anionic charges on the cell membrane surfaces, thus triggering saddle-splay curvature, which is correlated with the permeation ability of CPPs. On the other hand, protonated Lys is unable to promote the curvature in lipidic membranes and consequent cell penetration.^{12, 16} Based on these evidences, guanidinium-rich synthetic analogues of the CPPs have been developed.^{12, 15} It has been reported that the structural conformation of CPPs is not a prerequisite for their cell penetration activity even though the conformation dictates their penetration mechanism. CPPs that possess random conformation, such as TAT or Arg9, undergo endocytic pathway rather than translocation, which is typical for CPPs with alfa-helix and beta-strand conformations. Thus, the absence of a specific structural conformation of this class of cell penetration systems may be preferred when they are exploited for the delivery of macromolecular and colloidal cargoes and thus appealing for biomedical applications.¹⁷

Despite CPPs have extraordinary potential in the development of high performance drug delivery systems, they suffer from cell non-selectivity, thus yielding unspecific tissue association and random biodistribution, which represents a major drawback for their biomedical application for therapeutic purposes. Therefore, the use of CPPs for cell targeting requires dedicated molecular strategies to provide drug nanocarriers with site-selectivity. Furthermore, the use of these

polypeptides is limited by their poor stability in biological environment due to proteolytic degradation.¹⁸⁻¹⁹ To overcome these limitations, we developed a stable novel non-peptide Arg based dendritic cell penetrating enhancer (CPE), tetraArg-[G-1]-distearoyl glycerol (Arg₄-DAG) for liposomes decoration. Arg₄-DAG contains a non-symmetrical dendritic scaffold based on 2,2-bis(hydroxyl-methyl)propionic acid (bis-MPA) building block that, on one side, terminates with oligo-cationic Arg with higher charge density than that found in natural linear CPPs, and, on the opposite side, with a hydrophobic tail to anchor the CPE to the liposome bilayer. In order to enable the Arg₄-DAG decorated liposomes with tumor targeting behavior, the cationic charges were charge-to-charge quenched with a releasable PEGylating agent, mPEG_{5kDa}-oligomethacriloyl sulfadimethoxine (mPEG_{5kDa}-SDM₈), which includes a pH-sensitive poly-anionic block. The mPEG_{5kDa}-SDM₈ coated Arg₄-DAG decorated liposomes have been designed to bestow pH-switchable liposomes. Under physiological conditions at pH 7.4, namely in blood, the liposomes possess *stealth* features that minimize the association of the carrier to healthy cells. Under the acid environment of the tumor,^{4, 20} the oligo-Arg moieties on the liposome surface are revealed upon mPEG_{5kDa}-SDM₈ detachment thus enhancing the association of the liposomes to cancer cells.

2. EXPERIMENTAL SECTION

2.1 Materials and equipment

Chemicals and equipment are reported in the electronic supplementary information (ESI).

First (G1) and second (G2) generation bis-MPA dendrons bearing 4 and 8 Arg, respectively, and functionalized with diacyl glycerol, were designed *in silico* by ChemDraw (PerkinElmer Inc. Waltham, MA, USA). The structures underwent energy minimization using Force MMFF94x by Molecular Operating Environment (MOE) software (Chemical Computing Group Inc. Montreal, Canada). The data were elaborated assuming Arg monomers as dots centred on guanidinium carbon.

The reaction scheme illustrates the synthesis of cell-penetrating peptides (CPPs) 10 and 14. The synthesis begins with the conversion of 2,2-bis(hydroxymethyl)propionic acid (bis-MPA) to 1 (80% yield) and then to 2 (80% yield). 2 is converted to 3 (85% yield) and then to 4 (90% yield). 4 is converted to 5 (78% yield) and then to 6 (quantitative yield). 6 is converted to 7 (71% yield) and then to 8 (92% yield). 8 is converted to 9 (71% yield). 9 is converted to 10 (83% yield). 10 is converted to 14 (95% yield). The scheme also shows the synthesis of 11 (97% yield) and 12 (82% yield). 12 is converted to 13 (95% yield). 13 is converted to 14 (95% yield). The legend indicates: red line = 2,2-bis(hydroxymethyl)propionic acid scaffold (bis-MPA); green line = 2,2-bis(aminomethyl)propionic acid scaffold (bis-MPA); purple line = 3-(1,2,3-triazol)propane-1,2-distearate anchoring unit; blue line = arginyl cell penetration enhancer unit.

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^aReagents and conditions: (i) BnBr, KOH, DMF, 100 °C, 15 h; (ii) TsCl, Py, DMAP, CH₂Cl₂, 60 °C, 4 h; (iii) NaN₃, DMF, 90 °C, 16 h; (iv) 1) PtO₂, H₂, (Boc)₂O, MeOH, rt, 4 h; 2) 10% Pd/C, H₂, MeOH, rt, 3 h (v) DMP, PTSA, Acetone, rt, 2 h; (vi) DCC, CH₂Cl₂, rt, 4 h; (vii) 2-propyn-1-ol, Py, DMAP, CH₂Cl₂, rt, 15 h; (viii) Dowex 50W-X8 (H⁺-form), MeOH, rt, 6 h; (ix) 1) **4**, DCC, CH₂Cl₂, rt, 16 h; 2) **8**, Py, DMAP, CH₂Cl₂, rt, 16 h; (x) 1) CH₂Cl₂/TFA (1:1), rt, 2.5 h; 2) Boc-Arg(Pbf)-OH, HOBt, EDC, DIPEA, DMF, rt, 16 h; (xi) stearoyl chloride, Py, CH₂Cl₂, rt, 16 h; (xii) NaN₃, DMF, 100 °C, 15 h; (xiii) CuI (10 mol %), AcOH (20 mol %), DIPEA (20 mol %), sodium ascorbate (10 mol %), CH₂Cl₂, rt, 24 h; (xiv) 1) CH₂Cl₂/TFA (2.5:1), rt, 24 h; 2) CH₃CN/H₂O + 0.05% TFA (1:1), Amberlite IRA-900 (Cl⁻-form).

Briefly, compound (**14**) was synthesized by the Cu(I)-catalyzed Huisgen 1,3-dipolar cycloaddition (“click” reaction) between the azide terminated distearoyl propane anchoring unit (**12**) and the alkynic dendron (**10**) to give the Pbf-Boc-protected precursor (**13**), followed by TFA mediated deprotection of the Arg units (**xiii** and **xiv**, Scheme 1). The 3-azidopropane-1,2-distearate (**12**) was prepared by a two-step high yield procedure from commercially available (±)-3-chloro-1,2-propanediol (**3-MCPD**): acylation with stearoyl chloride followed by nucleophilic substitution of chloride with azide (**xi** and **xii**, Scheme 1). The synthesis of the four branched alkynic tetra-Boc-arginyl(Pbf) scaffold (**10**) was achieved via anhydride coupling between 2,2-(bis((*tert*-butoxycarbonyl)amino)methyl)-propanoic acid (**4**) and propargyl 2,2-bis(hydroxymethyl)propanoate (**8**) to yield (**9**) (**ix**, Scheme 1), followed by TFA mediated Boc deprotection and coupling with Boc-Arg(Pbf)-OH (**x**, Scheme 1). Compound (**8**) was synthesized according the procedure reported by Barnard et al. with only slight modifications²¹ (**v** - **viii**, Scheme 1). Compound (**4**) was synthesized by benzylation of the carboxylic function of bis-MPA followed by tosylation of the two free hydroxyl groups and nucleophilic substitution of the tosylate with azide to obtain (**3**) which was then reduced in a one-pot hydrogenation procedure

involving two sequential steps: the first in the presence of PtO_2 and Boc-anhydride to obtain the corresponding Boc-protected diamine and the second in the presence of Pd/C to cleave the benzyl protecting group and obtain the desired intermediate (**4**) (**i** - **iv**, Scheme 1).

2.3 Synthesis and characterization of the pH-sensitive copolymer mPEG-(poly-methacryloyl sulfadimethoxine) (mPEG_{5kDa}-SDM₈)

mPEG-polySDM (**17**) was synthesized in a two-step procedure, according to a modified strategy reported in the literature.²² The activated mPEG_{5 kDa}-NH-CO-C-(CH₃)₂-Br (**16**) was obtained by α -bromoisobutyryl bromide reaction with mPEG_{5kDa}-NH₂ (yield 90%). The TNBS spectrophotometric analysis confirmed the complete functionalisation of the amino groups of mPEG_{5kDa}-NH₂ with α -bromoisobutyryl bromide. The integrals of the ¹H NMR signals (δ 3.65 for the oxyethylene monomers of PEG and δ 1.9 for the two methyl groups of the α -bromoisobutyryl amide) were in agreement with a 1:1 PEG/ α -bromoisobutyryl amide molar ratio of the AGET-ATRP initiator. The synthesis of mPEG-polySDM (**17**) was performed by AGET-ATRP polymerization of methacryloyl sulfadimethoxine (SDM, (**15**)) using a 1:12 mPEG_{5 kDa}-NH-CO-C-(CH₃)₂-Br/SDM molar ratio. The polymer was isolated with a 67% yield. The ¹H NMR integrals (Figure S1 in ESI) of the signals at 6.0 ppm and 3.5 ppm, corresponding to the pyrimidyl hydrogen of SDM and the PEG oxyethylene monomers, respectively, showed that the co-polymer included, on average, 8 methacryloyl sulfadimethoxine monomers which was confirmed by UV-Vis spectroscopic analysis and iodine assay.

2.4 Liposome preparation

Liposomes were prepared according to the “thin layer rehydration technique” method,²³ using a 2:1 mol/mol hydrogenated soy phosphatidylcholine/cholesterol mixture. Briefly, 10 mg of the lipid mixture was dissolved in 1.5 mL of CH₂Cl₂ and the organic solvent was removed under reduced pressure by rotavapor to form a lipid film and then overnight to remove any CH₂Cl₂ trace. The lipid film was rehydrated with 200 μ L of buffer (0.01 M phosphate, 0.15 M NaCl at pH 7.4 or 0.01 M HEPES, 0.15 M NaCl at pH 7.4 and processed with ten freeze-thawing cycles.

CPE-decorated liposomes were prepared by adding 0.5 mL liposome dispersions (10 mg/mL lipid concentration) of increasing volumes of a 5 mg/mL Arg₄-DAG solution in the same buffer of the liposome dispersion in order to yield Arg₄-DAG in the range of 1-8 mol% with respect to lipids. The liposomes were sonicated for 60 sec with a Ultrasonic Homogenizer (Omni International, Kennesaw, GA-USA) set at 20% power. Non-associated CPE was removed by liposome centrifugation for 30 min at 11,000 rpm. The liposomes were redispersed with the same buffer to a lipid concentration of 5 mg/mL and extruded eleven times at 60 °C through a 200-nm cut-off polycarbonate membrane. The formulations were then incubated at 37 °C for 1 hour and the Arg₄-DAG content was assessed by Sakaguchi assay as reported in the ESI.

Liposomes decorated with 4 mol% of Arg₄-DAG with respect to lipids were incubated with increasing amounts of mPEG_{5kDa}-SDM₈ (0-4 mol% mPEG_{5kDa}-SDM₈/lipids molar ratio) in the same buffer to obtain PEG coated pH-sensitive formulations and diluted 50 times either with 0.01 M HEPES, 0.15 M NaCl at pH 7.4 and 7.0, or with 0.01 mM MES, 0.15 M NaCl at pH 6.5 and 6.0 and the zeta potential was assessed as a proxy for the Arg₄-DAG shielding and unshielding, respectively.

Liposomes decorated with 4 mol% of mPEG_{5kDa}-DSPE with respect to lipids were prepared as control with non-pH releasable PEG using a 5 mg/mL stock solution of mPEG_{5kDa}-DSPE in the same buffers.

Stability of liposomes over time was assessed at predefined time points by size and zeta potential analysis.

Fluorescent liposomes were prepared by including 0.2 mol% Rhodamine-DHPE with respect to lipids in the lipid film.

2.5 Particle size and zeta potential measurements

Size and polydispersity, and zeta potential of liposomes were assessed by *Dynamic Light Scattering* (DLS) using a NanoZS Zetasizer (Malvern, Worcestershire, UK). The analyses were performed by using liposome dispersions obtained by dilution as reported above in different buffers: 0.01 M phosphate, 0.15 M NaCl (PBS) at pH 7.4 and 6.5; 0.01 M phosphate containing 300 mM mannitol (PBS/mannitol) at pH 7.4 and 6.5; 0.01 M HEPES, 0.15 M NaCl (HEPES) at pH 7.4; 0.01 M HEPES containing 300 mM mannitol (HEPES/mannitol) at pH 7.4; 0.01 M MES, 0.15 M NaCl (MES) at pH 6.5; 0.01 M MES containing 300 mM mannitol (MES/mannitol) at pH 6.5. The liposome size was expressed as z-average.

Zeta potential analyses were performed using liposomes decorated with 4 mol% of Arg₄-DAG and coated with increasing of mPEG_{5kDa}-SDM₈/lipids molar ratio in HEPES at pH 7.4 and MES at 6.5.

2.6 SAXS Analysis

The internal structure of liposomes was analyzed by synchrotron small-angle X-ray scattering (SAXS) using an ID02 high-brilliance beamline (ESRF, Grenoble, France), with a beam cross section $200 \times 400 \mu\text{m}$ and wavelength $\lambda = 0.1 \text{ nm}$, investigated momentum transfer region, $q = (4\pi/\lambda) \sin(\theta)$, $0.0116 < q < 40 \text{ nm}^{-1}$, 2θ scattering angle.²⁴⁻²⁵ The corresponding real-space length-scale was from 100 to 0.1 nm. Liposome (10 mg/mL lipid concentration), CPE (5 mg/mL) and mPEG_{5kDa}-SDM₈ (5 mg/mL) solutions in HEPES at pH 7.4 and MES at 6.5 were transferred in plastic capillaries (KI-BEAM; ENKI srl, Concesio, Italy), 2 mm internal diameter. Measurements were performed at 25 °C and the data were treated for angular regrouping and background subtraction, and then reported as scattered radiation intensity as a function of the momentum transfer q .

2.7 MP-SPR measurements

Surface plasmon resonance measurements were performed with a multi-parametric MP-SPR Navi™ 200 (BioNavis Ltd., Tampere, Finland) instrument. The system was equipped with two temperature controlled independent flow channels, inlet and outlet (waste) tubings, and a manual sample injector for both flow channels. The SPR measurements were performed with a laser wavelength of 670 nm. The sensor chip temperature was set to 20 °C. CPE-decorated liposomes were captured and immobilized on SPR sensors consisting of a thin 6-kD carboxymethyl dextran hydrogel layer functionalized with dodecyl lipid anchors.²⁶ The pH-sensitive polymer (mPEG_{5kDa}-SDM₈) was dissolved in running buffers (HEPES, pH 7.4, and MES, pH 6.5) to

assess polymer/liposome association under different pH conditions. The flow rates used for immobilizing the CPE-decorated liposomes and for monitoring the interactions between the pH-sensitive polymer (mPEG_{5kDa}-SDM₈) and immobilized liposomes were 50 and 100 $\mu\text{L}/\text{min}$, respectively.

The SPR sensors were used repeatedly after rejuvenation with an injection series of Hellmanex II 2% or CHAPS 20 mM, ethanol 80% and Milli-Q water. In between measurements, the sensors were stored immersed in CHAPS at 4 °C. During the SPR measurements, the functionalized gold sensor slides were first subjected to the running buffer for the respective liposomes for approximately 5-10 min until a stable baseline was achieved. In the second phase, the liposomes were injected into both flow channels for 25 min, and lastly, the sensor surfaces with the immobilized liposomes were exposed to seven sequential injections of 0.4 μM , 0.8 μM , 1.61 μM , 3.23 μM , 6.45 μM and 12.9 μM pH-sensitive polymer (mPEG_{5kDa}-SDM₈) solutions in the buffers at pH 7.4 or 6.5.

2.8 Liposome loading and release study

Liposome loading with BSA-Rhodamine (BSA-Rho) and calcein and the corresponding method of the release study are reported in the ESI.

2.9 Cell cultures

HeLa cells (human cervical cancer) were grown at 37 °C in DMEM medium supplemented with 10% (v/v) heat-inactivated fetal bovine serum (FBS), 2 mM L-glutamine, 100 IU/mL penicillin, 100 µg/mL streptomycin and 0.25 µg/mL of amphotericin B, with a 5% CO₂ atmosphere. The cells were harvested by treatment with 0.05% (w/v) trypsin-0.02% (w/v) EDTA solution (Sigma-Aldrich, St. Louis, MO-USA), suspended in culture medium and seeded at the appropriate cell concentration.

Viability studies were performed by incubation of HeLa cells with the different liposome formulations according to the procedure reported in the ESI.

2.10 Cytofluorimetric analysis

HeLa cells were used to evaluate the internalization of: naked liposomes (plain liposomes); PEGylated liposomes (liposomes coated with non-pH-responsive mPEG_{5kDa}-DSPE); liposomes decorated with 2 or 4 mol% of Arg₄-DAG; pH-responsive liposomes decorated with 4 mol% Arg₄-DAG and coated with 4 mol% of mPEG_{5kDa}-SDM₈, labelled with rhodamine-DHPE, or loaded with BSA-Rho or calcein. The cells were seeded in a 6 well plate at a density of 1×10⁶ cells/well. After 24 hours, the cells were washed twice with HEPES or MES at pH 7.4 and 6.5 respectively and incubated for 1 hour at 37 °C with 1 mL of 0.1 mg/mL liposomal dispersion in DMEM supplemented with 10% of FBS buffered with 0.1 M HEPES, pH 7.4, or 0.1 M MES, pH 6.5.²⁷ The medium was removed and the cells were washed three times with fresh PBS at pH 7.4 and treated for 2 min with 300 µL of 500 µg/mL trypsin in PBS without calcium and

magnesium. DMEM (1 mL) supplemented with 10% FBS was added to the wells to quench trypsin and the cells were recovered and centrifuged at 1,000 rpm for 5 minutes. The cellular pellet was resuspended in PBS, pH 7.4, washed twice with the same buffer and recovered by centrifugation at 1,000 rpm for 5 minutes. The cell samples were resuspended in 300 μ L of PBS pH 7.4, and analysed by flow cytometry using a BD FACScanto II flow cytometer (Biosciences, San Jose, Canada) equipped with a FACSDIVA software package. The cell populations were gated using forward versus side scatter to exclude debris and dead cells.

The same protocol was also used with PNT2 cells used as non-cancer cell model.

2.11 Confocal microscopy

The liposome association to cancer cells and cell uptake were studied by confocal microscopic imaging.

HeLa cells were seeded onto 4-chamber tissue culture microscope slides at a 1×10^5 cells/chamber density. After 24 hours (90% confluence), the cells were washed twice with HEPES at pH 7.4 or MES at pH 6.5, and the medium replaced with 0.5 mL of 0.1 mg/mL liposome dispersions in DMEM supplemented with 10% of FBS buffered with 0.1 M HEPES, pH 7.4, or 0.1 M MES, pH 6.5:²⁷ Rhodamine-DHPE labelled naked liposomes (plain liposomes), liposomes coated with non-pH-responsive mPEG_{5kDa}-DSPE (PEG/liposomes), liposomes decorated with 4 mol% Arg₄-DAG (CPE/liposomes), liposomes decorated with 4 mol% Arg₄-DAG and coated with 4 mol% mPEG_{5kDa}-SDM₈ liposomes (CPE/pH-sensitive polymer / liposomes). Liposomes loaded with BSA-Rho or calcein were also tested. After

incubation for 1 hour at 37 °C, the medium was removed and the cells were gently washed three times with PBS, pH 7.4, and finally fixed by treatment with 4% paraformaldehyde in PBS, pH 7.4, for 10 min in an ice bath. The cell membranes were stained by cell incubation with 5 µg/mL Alexa Fluor® 633 conjugated wheat-germ agglutinin (WGA-AlexaFluor 633) in PBS, pH 7.4, for 10 minutes at room temperature. The cells were then washed with PBS, pH 7.4, and the samples were covered with a glass slide using Vectashield® as mounting medium containing 1.5 µg/mL DAPI for nucleus staining. The samples were maintained at 4 °C in the dark until microscopic examination.

The samples were imaged by confocal laser microscopy using a LSM 800 microscope (Carl Zeiss, Jena, Germany) equipped with a 63x, n.a. 1.4, oil immersion objective and a ZEN 2.1 - blue edition - software (Carl Zeiss, Jena, Germany). Lasers with emission wavelength at 405, 488, 561 and 640 nm were used to detect DAPI, calcein, Rhodamine-DHPE and WGA-Alexa Fluor 633, respectively. To avoid emission crosstalk, each emission fluorescence was recorded independently with specific detector and optical cut-off filter over the entire emission spectrum of related chromophores. Image analyses were performed using ImageJ 1.47v (National Institutes of Health software package).

2.12 Statistical analysis

Statistical analyses were performed with XLSTAT software (New York, USA). Two-way analysis of variance was used to calculate the threshold of significance. Statistical significance was set at $p < 0.05$.

3. RESULTS AND DISCUSSION

The work presented here was aimed at developing a new novel strategy that provides for controlled cell up-take of nanocarriers by exploiting environmentally triggerable cell penetration enhancers (CPEs) and reversible *stealth* polymers. The nanocarrier has been designed as an alternative to systems that exploit conventional “active targeting” based on cell biorecognition. Indeed, this nanocarrier can provide for intracellular drug delivery after local microenvironmental activation.

Although many efforts have been done to exploit CPEs, namely TAT derived peptides, to ameliorate the therapeutic performance of drug nanocarriers, non-specificity and lack of selective targeting strongly limits their use for fabrication of smart nanomedicines.²⁸⁻³⁰ Therefore, CPE exposure control by environmental stimuli is required to generate drug nanovehicles for site-selective cell targeting.

Nanocarrier *stealthiness* is a requisite to prevent their clearance by RES. On the other hand, *stealthiness* may reduce the interaction of colloidal drug carriers with the target cancer cells, which may result in poor access to the cytosol, low intracellular drug delivery and, eventually, reduced therapeutic performance.

Thus, we synthesised a new non-peptide, non-linear CPE and a pH-sensitive *stealth* block-copolymer, which were synergistically combined to produce “smart” liposomes that undergo a four-stage targeting process: 1. circulate in the bloodstream in a dormant conformation; 2. passively dispose in the tumor according to the enhanced permeability and retention (EPR)

mechanism;³¹⁻³² 3. are activated by microenvironmentally controlled CPE exposure; 4. are taken-up by the cells.

3.1 Dendron cell penetration enhancer (CPE) design, synthesis and characterization.

The novel CPE formed by multiple Arg units to the tips of a first-generation dendritic structure was designed in order to develop a molecule with enhanced physicochemical and biological features compared to synthetic and natural TAT-like peptides (CPPs), namely high stability towards proteases, lack of bioactivity and high biocompatibility. Furthermore, Sheldon reported that cell penetrating peptides with dendritic architecture typically outperform their linear monomeric homologues in term of cellular uptake.³³⁻³⁴

Multiple Arg units were conjugated to a first-generation dendritic structure based on a bis(aminomethyl)propionic acid (bis-AMPA) branched scaffold anchored to a core of 2,2-bis(hydroxymethyl)propionic acid (bis-MPA). This dendritic architecture was selected for the excellent biocompatibility, stability under physiological conditions and biodegradability of the bis-MPA core³⁵, which makes it suitable for clinical applications. Arg was coupled to the dendron through stable amidic bonds. Wender demonstrated that the chirality of the amino acids of TAT has a negligible effect on the cellular uptake of this class of molecules.³⁶ Therefore, the stereochemistry of the resulting Arg-decorated dendron was not an issue in this work.

The divergent synthesis of the bis-MPA/bis-AMPA based dendron yielded multiple-arm derivatives classified as generation (G) that corresponds to the consecutive branching points each of them end-terminating with four amino groups for Arg coupling. Notably, TAT possesses 6

Arg (sequence: GRKKRRQRRRPQ). Therefore, the G1 and G2 bis-MPA based dendrons, which bear 4 and 8 Arg, respectively, have the closest number of Arg to the natural TAT.

The charge density of CPPs has been shown to play a key role in the cell glycolyx cross-linking and cell penetration. Accordingly, we investigated the charge density of the G1 and G2 Arg-decorated dendrons as predicted from the average mutual distance of Arg using the Molecular Operating Environment (MOE) software (Chart 1).¹⁴

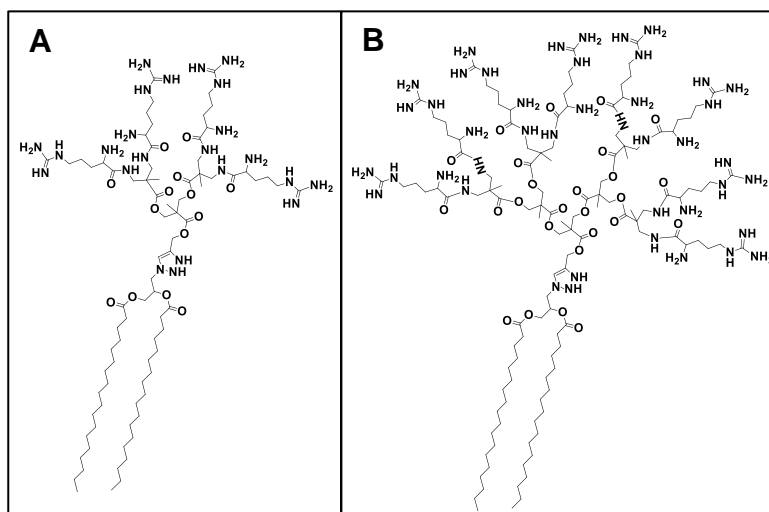


Chart 1. Structures of first generation (G1) dendron Arg₄-DAG (A) and second generation (G2) dendron Arg₈-DAG (B).

The *in silico* simulation showed that the carbon atoms of adjacent guanidinium Arg residues are separated by 10.3 Å and 13.8 Å in G1 (4 Arg) and G2 (8 Arg) dendrons, respectively. This result suggests that G1 dendron has higher charge density than G2 dendron. It is worth to note that You-Rim et al. investigated the correlation between the number of Arg and the cell penetration

performance of short peptides in nanocapsular systems, showing that carriers functionalized with short peptides with only three Arg are efficiently taken-up by the cells. This was ascribed to the cooperative effect that oligo-Arg on the nanocarrier surface.³⁷

The dendron design included the end-functionalization with a diacyl glycerol unit, 3-(1,2,3-triazol)propane-1,2-distearate (DAG), to yield its association with the liposomal membrane. Considering that the resulting [hydrophobic alkyl chains]/[polycationic dendritic block] mass ratio was 1:2.8 and 1:4.6 for the G1 derivative (4 Arg) and the G2 derivative (8 Arg), respectively, the former, that contains a heavier anchoring fraction with respect to the latter, was expected to more stably associate with the lipid bilayer of the liposome membrane. Therefore, taking into account the higher positive charge density and the lighter cationic block, the G1 dendron bearing four Arg (Arg₄-DAG) was selected.

The synthetic procedure set-up for the production of Arg₄-DAG provided for high structural control and monodispersity of the branched derivative, and high functional group density at the terminal ends (namely the Arg). Feliu et al.³⁵ investigated the stability, degradability and cytotoxicity of bis-MPA based dendrons under different conditions and demonstrated the excellent biocompatibility and potential for exploiting these scaffolds for biomedical applications.

3.2 pH-sensitive block-copolymer synthesis and characterization

The acid-sensitive mPEG-(poly-methacryloyl-sulfadimethoxine) (mPEG-polySDM) *stealth* block co-polymer (**17**) was designed to yield Arg₄-DAG shielding and unshielding at the surface of CPE-decorated liposomes.

The ionizable sulfadimethoxine (SDM) block was selected to provide the coulombic interaction of the co-polymer with the polycationic CPE. Actually, by virtue of its pKa (6.1),³⁸ SDM reversibly switches from anionic to uncharged state over a narrow pH range close to physiopathological conditions. According to the evidence that the pKa of SDM oligomers increases with the number of monomers,^{22, 39} a block mPEG-polySDM co-polymer formed by 7 SDM monomers was synthesised. The potentiometric titration profile reported in Figure S2 (ESI) shows that this copolymer has pKa 7.1. This result is in agreement with polymers obtained with PEG derivatives and SDM reported in the literature.^{22, 39-40} Therefore, this co-polymer is predominantly in the polyanionic form at pH 7.4 of blood, which allows for the charge-to-charge association with the CPEs on the liposome surface, while it switches to a less charged form at pH 6.5 of the tumor²² which translates in a weaker electrostatic interaction with the cationic liposomes resulting in detachment from their surface and CPE unmasking (see chapter 3.4).

The length of the PEG block was selected according to the literature reports showing 5 kDa mPEG minimizes the protein opsonisation of nanocarriers and their removal by the RES system, yielding prolonged permanence in the bloodstream.^{32, 41}

3.3 pH-sensitive liposome formulation

Arg₄-DAG decorated liposomes were prepared by rehydration of a thin lipid layer²³ and CPE decoration by a spontaneous micelle transfer post-insertion technique,⁴² which provided for the CPE exposure on the outmost layer of the lipidic vesicles and reduced the hiding of the Arg₄-DAG in the liposome core.

Liposomes decorated with increasing Arg₄-DAG/lipids molar ratio (from 1 to 8 mol%) were prepared according to the protocols reported in the literature for liposomes decorated with TAT-like derivatives.⁴³⁻⁴⁵ The Arg₄-DAG association to the liposome bilayer was assessed by zeta potential analysis. In PBS, pH 7.4, the non-decorated control liposomes exhibited a zeta potential that was nearly neutral (i.e. -1 mV) while the results reported in Figure 1A show that the zeta potential of the liposomes increased as the Arg₄-DAG concentration on the vesicle surfaces increased up to a plateau of +24 mV at 4 mol% Arg₄-DAG/lipid ratio. These results suggest that there is a limit of Arg₄-DAG association with the liposome membrane that could be explained with a charge repulsion effect, being the Bjerrum length at ~150 mM ionic strength still similar to that in water, ~7 Å. The limited Arg₄-DAG association was confirmed by the Sakaguchi assay,⁴⁶ which showed that more than 90% of the Arg₄-DAG was associated to the lipid bilayer when Arg₄-DAG/lipid molar ratios between 1 and 4 mol% were used to assembled liposomes while the efficiency of the CPE association to the lipid bilayer decreased from 90% to ~50% with 8% Arg₄-DAG/lipid molar ratio.

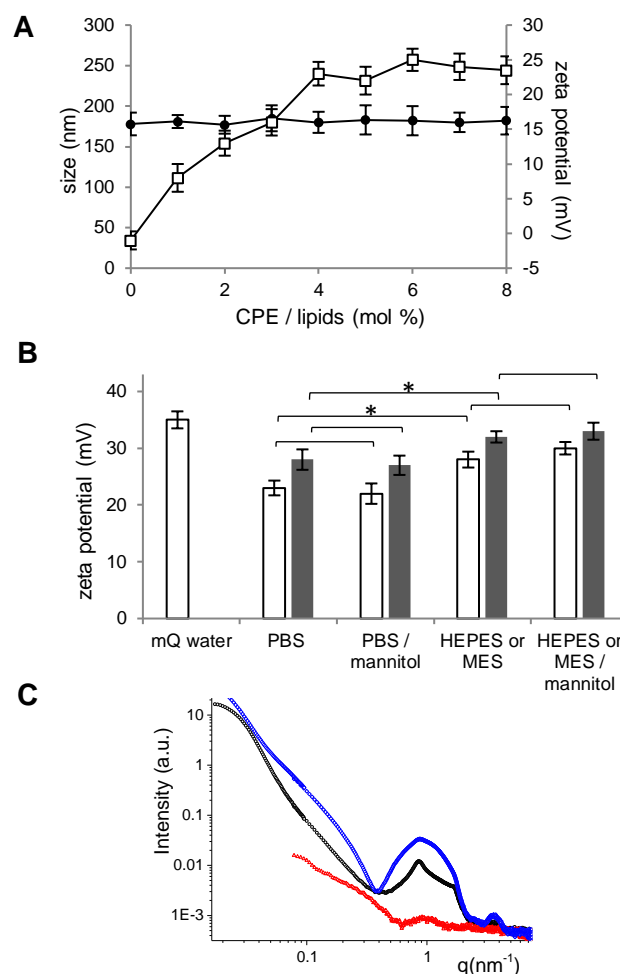


Figure 1. (A) Size (●) and zeta potential (□) profiles of liposomes decorated with increasing Arg₄-DAG/lipid ratio at pH 7.4 in PBS; (B) Zeta potential of liposomes coated with 4 mol% CPE at pH 7.4 (□) and 6.5 (■) in different buffers; statistical analysis: * p<0.05; ** p<0.01; *** p<0.001. (C) Small angle x-ray scattering spectra of CPE alone (■) and liposomes before (■) and after (■) incubation with Arg₄-DAG. T = 25 °C, HEPES buffer pH 7.4.

Zeta potential analyses were performed in HEPES/NaCl or HEPES/mannitol at pH 7.4 and in MES/NaCl or MES/mannitol at pH 6.5 to examine the effect of pH, buffer ionic strength and

osmotic activity on biophysical properties of Arg₄-DAG decorated liposomes.⁴⁷ The zeta potential values determined at pH 7.4 (mimicking the blood) were slightly lower than that at pH 6.5 (mimicking the tumour) regardless the buffers used for the analysis (Figure 1B). This result indicates that the overall positive charge on the liposome surface conveyed by the CPE increased as the pH decreased, despite the high pKa of the guanidinium group of Arg (pKa 12.48).⁴⁸ The positive charges of the Arg₄-DAG decorated liposomes in phosphate (anionic buffer) at both pHs (7.4 and 6.5) was slightly lower than in HEPES or MES (zwitterionic buffers) indicating that the anionic species of the buffer partially quenched the Arg of Arg₄-DAG. No significant differences were instead observed with different osmotic agents (mannitol and NaCl). Thus, charge shielding of Arg₄-DAG by buffer and osmotic agents in the aqueous medium may be related to the size of the buffer species and their overall molecular charge distribution.

Based on the zeta potential analysis, HEPES (at pH 7.4) and MES (at pH 6.5) were used as buffers for the extensive biophysical characterization of the pH-responsive liposomes under environmental pHs that mimic the healthy tissue and the tumor. These two buffers are extensively used to prepare and characterize colloidal drug carriers.⁴⁹

Figure 1A shows that the plain and the decorated liposomes had a vesicle size of 178 ± 14.5 nm and 180 ± 13.3 nm, respectively, with a narrow polydispersity index (PDI 0.06), which indicates that the decoration with the synthetic CPE did not alter the formulation size. Furthermore, no size alteration was observed after 48 hours incubation at 37 °C, which confirms a good colloidal stability of the vesicles as a result of the charge mediated repulsion that inhibited aggregation.

SAXS studies were performed to assess the structure of decorated liposomes. The X-ray scattering spectra reported in Figure 1C show that Arg₄-DAG (CPE) alone formed small micelles

(red squares in Figure 1C), which was confirmed by a fluorescence assay used to assess the critical micelle concentration (CMC= 40 μM). The 2:1 (mol/mol) phosphatidylcholine/cholesterol liposomes before and after incubation with Arg₄-DAG (black and blue dots) showed the fingerprint of the multi-layer vesicles. Before Arg₄-DAG incubation (black dots), two peaks with different intensity, at $q_1 = 0.86 \text{ nm}^{-1}$ and $q_2 = 1.72 \text{ nm}^{-1}$, were superimposed to the form factor of the individual lipid bilayer. The 1-2 periodicity of these peaks is typical of a stratified liposome, with a repetition distance $d = 2\pi/q_1 = 7.3 \text{ nm}$, which is in agreement with values reported in the literature for similar 2:1 (mol/mol) phosphatidylcholine/cholesterol layers.⁵⁰ The spectrum obtained with liposomes after incubation with CPE at pH 7.4 is similar to that obtained at pH 6.5 (Figure 2C, green diamonds). Upon incubation of the liposomes with the CPE, the intensity profile dramatically changed, indicating that interaction between admixed components occurred. The final spectrum could not be obtained by simple superposition of the two, as shown in Figure S3 of the ESI. The intensity enhancement of the single-bilayer form factor and its better definition was typical for charge-bearing liposomes, as compared to uncharged. Multilamellarity was still observed, with the same repetition distance, showing that CPE insertion occurred only in the external bilayer of the liposome. Then, the liposome constituted a support for the CPE that decorated its surface without deeply affecting the liposome structure.

3.4 pH-controlled shielding of CPE decorated liposomes

The cancer cell interaction were investigated using 4 mol% Arg₄-DAG decorated liposomes coated with the pH-sensitive block co-polymer mPEG_{5kDa}-SDM₈.

The zeta potential analysis showed that the increase of the [pH-sensitive polymer]/[Arg₄-DAG] molar ratio decreased the vesicle charge due to the coulombic interaction of the polyanionic co-polymer with the polycationic liposomes. The zeta potential values reported in Figure 2A show that at both pH, 7.4 and 6.5, the overall charge of the decorated liposomes decreased as the mPEG_{5kDa}-SDM₈/Arg₄-DAG molar ratio increased. At pH 7.4, the 1:1 mPEG_{5kDa}-SDM₈/Arg₄-DAG molar ratio yielded complete liposome charge quenching, while at pH 6.5 a residual charge (+11 mV) was registered. This behaviour is ascribable to the pH dependent ionization of the polyanionic co-polymer. At pH 7.4 and 6.5, an average of 5.2 and 1.6 sulfadimethoxines (SDM), respectively, was calculated to be in the anionic form for each mPEG_{5kDa}-SDM₈ unit (pK_a 7.1), which resulted in an unlike ability to quench the positive charges of the Arg of the CPE and thus associating with the CPE-decorated liposomes. Notably, since the pH-sensitive liposomes at pH 6.5 are generated by pH shift from formulations at pH 7.4, the positive zeta potential of these liposomes at pH 6.5 supports a significant detachment of the mPEG_{5kDa}-SDM₈ during pH shifting as a consequence of the decrease of the co-polymer anionic charges (as discussed in chapter 3.2). On the contrary, the zeta potential is nearly neutral when pH-sensitive liposomes are generated at pH 7.4. The residual anionic charge of the polymer at pH 6.5 due to its apparent pK_a may account for the incomplete recovery of the zeta potential of liposomes at this pH since a fraction of the polymer may be still associated to the vesicles surface. Indeed, the detachment of mPEG_{5kDa}-SDM₈ from liposome surface started at pH below 7 and was more remarkable as the pH of incubation decreased to 6.0 as a result of more extensive polymer detachment from liposomes (Figure S4).

Zeta-potential analysis of Arg₄-DAG decorated liposomes incubated with 1:1 mPEG_{5kDa}-OH/Arg₄-DAG molar ratio showed that the ionically neutral 5 kDa mPEG_{5kDa}-OH was not able

to associate with Arg₄-DAG. Similarly to other anionic species present in the saline buffer, also the ionisable single SDM molecules incubated with the Arg₄-DAG decorated liposomes at a 8:1 SDM/Arg₄-DAG molar ratio were not found to associate with the Arg₄-DAG indicating that the cooperative effect of multiple anionic charges of the oligo-ionic mPEG_{5kDa}-SDM₈ was required to associate with the CPE. Notably, this electrostatic interaction finally resulted in the physical PEGylation of the liposomes at pH 7.4 and charge screening.

Size analysis showed that at pH 7.4 the mPEG_{5kDa}-SDM₈ coating of Arg₄-DAG decorated liposomes yields a slight vesicle size increase, which supports for the association of the pH-responsive polymer to the CPE decorated liposomes in agreement with data reported in the literature for post-insertion PEGylation of liposomes with 5 kDa PEG.⁵¹ The size and zeta potential analyses showed that the polymer coated vesicles were colloidally stable over 2 days while CPE-decorated liposomes (no mPEG_{5kDa}-SDM₈) showed instability after 48 hours (Figure S5 in ESI).

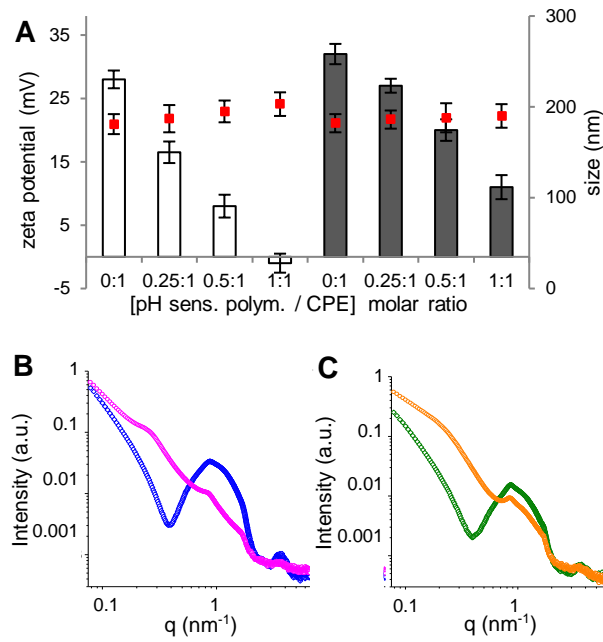


Figure 2. (A) Zeta potential (bars) and size (■) of Arg₄-DAG decorated liposomes in the presence of mPEG_{5kDa}-SDM₈ at increasing mPEG_{5kDa}-SDM₈/Arg₄-DAG (pH sens. polym. / CPE) feed molar ratio, at pH 7.4 (□), 6.5 (■). (B) Small angle x-ray scattering spectra of Arg₄-DAG decorated at pH 7.4, before (♦) and in the presence of mPEG_{5kDa}-SDM₈ (♦) and (C) at pH 6.5, before (♦) and in the presence of mPEG_{5kDa}-SDM₈ (♦).

SAXS spectra of Arg₄-DAG decorated liposomes at pH 7.4 (B) and pH 6.5 (C) reported in Figure 2 show that the mPEG_{5kDa}-SDM₈ addition modified the scattering profile of the liposomal dispersions. The internal structure was maintained for both systems, as shown by the persistence of the intensity peak at $q_1 = 0.86 \text{ nm}^{-1}$, revealing that the multi-layered liposomal core structure was preserved, with the same repeat distance, 7.3 nm. Also, intensity oscillations in the higher q region of the spectra display similar features, indicating that the individual lamellae maintain the same thickness. However, it is evident that the scattering profile was more modified at pH 7.4

(Figure 2B) than at pH 6.5 (Figure 2C) by the addition of mPEG_{5kDa}-SDM₈. Altogether, these results suggest that modifications were mainly related to the structure of the outer shell of the liposomes. At pH 7.4 a large association of the pH-sensitive polymer with the external surface of decorated liposomes can be inferred, while at lower pH the scattering profile was consistent with the presence of free PEG in solution. An example of reconstruction of the experimental data is reported in Figure S6 of the ESI.

Surface plasmon resonance (SPR) studies were performed to further investigate the association of the pH-sensitive polymer with CPE decorated liposomes immobilized on sensorchip (Figure S7 in ESI).^{26, 52-53} The real-time SPR profiles to increasing additions of mPEG_{5kDa}-SDM₈ to Arg₄-DAG liposomes reported in Figure 3A show that at pH 7.4 the SPR response was considerably higher than at pH 6.5. This result confirms that mPEG_{5kDa}-SDM₈ interacted more efficiently at pH 7.4 than at pH 6.5 as a consequence of the fact that at pH 7.4 and pH 6.5 each mPEG_{5kDa}-SDM₈ unit possessed on average 5.2 and 1.6 SDM in the anionic state, respectively.

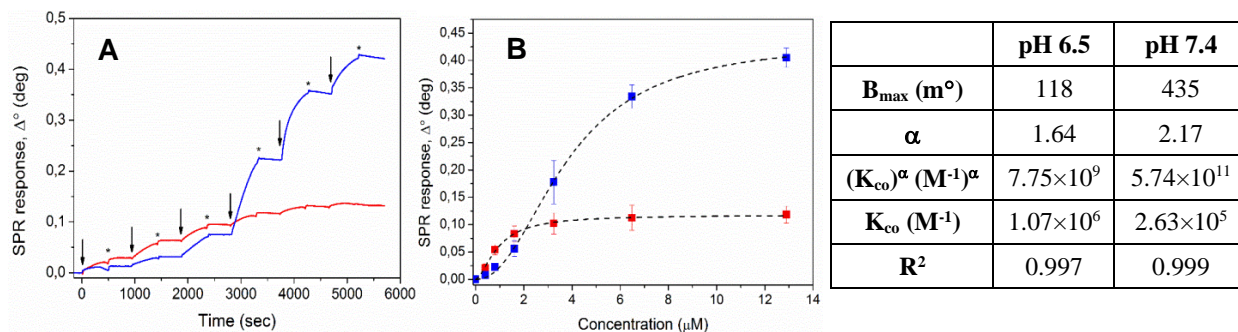


Figure 3. (A) Representative MP-SPR signal responses during interaction of increasing concentrations of mPEG_{5kDa}-SDM₈ at pH 6.5 (—) and 7.4 (—) through in situ PEGylation of Arg₄-DAG liposomes immobilized on the SPR sensor surface. Down arrows indicate injection time points of the pH-sensitive polymer, while stars indicate injection time points for the running

buffer. **(B)** Binding isotherms at pH 6.5 (■) and 7.4 (■) for mPEG_{5kDa}-SDM₈ interaction with Arg₄-DAG decorated liposomes immobilized on the SPR sensor surface (N=3). Dashed black lines are fits of the co-operative binding model to the data points. **Table:** B_{max} values, cooperativity coefficients (α), overall cooperative binding constant (K_{co}) ^{α} , and average cooperative binding constant (K_{co}) at pH 6.5 and 7.4 for mPEG_{5kDa}-SDM₈ interaction with Arg₄-DAG liposomes immobilized on the SPR sensor surface.

Since the association of mPEG_{5kDa}-SDM₈ with Arg₄-DAG liposomes involves multiple charge interactions, multivalent ligands and multi-subunit substrates, the binding constants and mechanism could be determined by a co-operative binding model analysis according to the following equation:⁵⁴⁻⁵⁵

$$B = \frac{[P]^\alpha (K_{co})^\alpha}{1 + [P]^\alpha (K_{co})^\alpha} \quad (1)$$

where B = SPR signal response, [P] = concentration of mPEG_{5kDa}-SDM₈, α = co-operativity coefficient and K_{co} = average cooperative binding constant. If the co-operativity of the binding is positive, then the first ligand facilitates the binding of the next ligand. If the co-operativity of the binding is negative, then each subsequent ligand is bound less strongly than the previous one. One-site, positive and negative co-operative binding models are obtained when the co-operativity coefficient α is 1, $\alpha > 1$ and $\alpha < 1$, respectively. The binding constants K_1, K_2, \dots, K_N for each charged interaction will not be the same and, therefore, only average co-operative binding constants can be obtained, i.e. K_{co} that is the average co-operative binding constant and $K_{tot} = (K_{co})^\alpha$ that is the overall average co-operative binding constant.

Figure 3B shows the co-operative binding isotherm of mPEG_{5kDa}-SDM₈ interaction with Arg₄-DAG decorated liposomes immobilized on the SPR sensor surface. The regression coefficients fitting co-operative models at pH 6.5 and 7.4 were 0.997 and 0.999, respectively. The B_{\max} value obtained at pH 7.4 was almost 4 times higher than at pH 6.5, which was consistent with much higher association of mPEG_{5kDa}-SDM₈ with the Arg₄-DAG decorated liposomes at pH 7.4 as discussed above. The co-operativity coefficients at both pH values are clearly larger than 1, which suggests that the interaction of mPEG_{5kDa}-SDM₈ with Arg₄-DAG liposomes displays a positive co-operativity at both pH values. The overall average binding constants $K_{\text{tot}} = (K_{\text{co}})^{\alpha}$ had very high values and was higher at pH 7.4 (i.e. 5.74×10^{11}) than at pH 6.5 (i.e. 7.75×10^9), which once again corroborates the higher association of mPEG_{5kDa}-SDM₈ with Arg₄-DAG liposomes at pH 7.4. However, the average binding constants (K_{co}) showed a reverse trend indicating a higher value at pH 6.5 (i.e. 1.07×10^6) compared with pH 7.4 (i.e. 2.63×10^5). This kind of behavior is expected for systems exhibiting strong positive co-operativity as the individual binding constant increases with each bound ligand, i.e. $K_1 < K_2 < K_3 \dots$ until the saturation limit is reached, which results in that $K_{\text{tot}} \gg K_{\text{co}}$. Furthermore, the smaller the K_{co} value the higher the charge ratio is required to reach the saturation level of the system, which was also reflected in the interaction of mPEG_{5kDa}-SDM₈ with Arg₄-DAG liposomes when comparing the two different pH conditions.

Since liposomes are exposed to serum proteins upon parenteral administration, stability studies were performed by incubation in foetal bovine serum (FBS) at pH 7.4 and 6.5 (Figure 4). The size increase of CPE-decorated liposomes at both pHs could be ascribed to a combined effect of “*protein corona*” formation due to the high positive zeta potential and possible aggregation. This hypothesis is in agreement with results reported in the literature⁵⁶⁻⁵⁷ including the SPR

measurements of protein adsorption on the oligo-arginyl decorated liposomes we reported in our previous studies.⁵³ mPEG_{5kDa}-SDM₈ coating reduced the liposome size increase at pH 7.4 indicating that the pH-sensitive polymer reduced the opsonization of the cationic liposomes. This result is also in agreement with our SPR outcomes reported in the literature showing that the CPE decorated liposomes coated with polyanionic-PEG display a thinner protein corona with respect to the polyanionic-PEG free formulation.⁵³ On the other hand, at pH 6.5 in the presence of FBS, the dissociation of the mPEG_{5kDa}-SDM₈ from the liposome induced a size increase of the vesicles that was comparable to that of the mPEG_{5kDa}-SDM₈ free CPE-decorated liposomes confirming that, under this pH condition, the liposome surface was prevalently exposed to protein adsorption.^{32, 58-59}

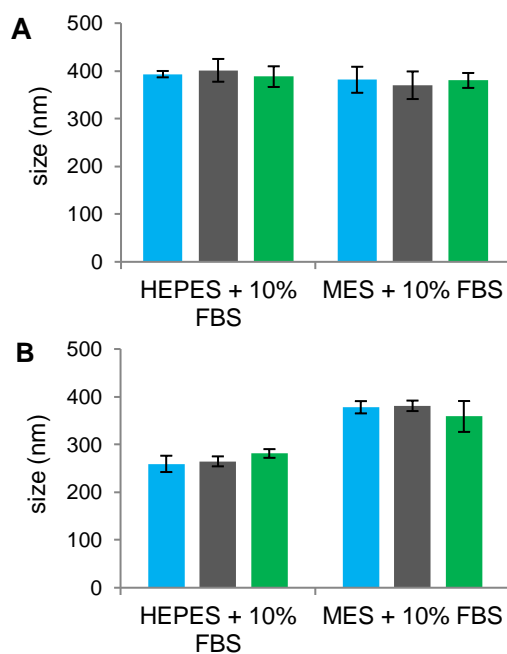


Figure 4. Size of CPE decorated liposomes (A) and CPE decorated liposomes formulated with mPEG_{5kDa}-SDM₈ (B) in 0.01 M HEPES, 0.15 M NaCl at pH 7.4 and 0.01 M MES, 0.15 M NaCl at 6.5 after 0 (■), 1 (■), 2 (■) hours from addition of 10% FBS.

3.5 Biological Studies

3.5.1 Effect of CPE on liposome association to cells

Cell association and internalization studies were carried out using naked liposomes (control) and 2 and 4 mol% Arg₄-DAG decorated liposomes labelled with rhodamine-DHPE.

The cytofluorimetric results reported in Figure 5A show that the cell association increased with the Arg₄-DAG density on liposome surface. The decoration of liposomes with 4 mol% CPE (zeta potential = +28 mV) increased the liposome association to cells of 31 times (referred to the MFI) with respect to naked liposomes (zeta potential = -1 mV) and 11 times with respect to 2 mol% Arg₄-DAG decorated liposomes (zeta potential = +13 mV).

Cell association of the CPE-decorated liposomes can be mediated by the protein corona adsorbed on liposome surface.⁵⁶ However, despite the presence of the protein corona, interaction of the CPEs with the cell glycocalyx when liposomes approach the cell membrane may take place and mediate the association of the liposomes with HeLa cells. Further investigations are required to elucidate in details at molecular level the mechanism of cell entry mediated by this novel CPE which is a quite complex event.

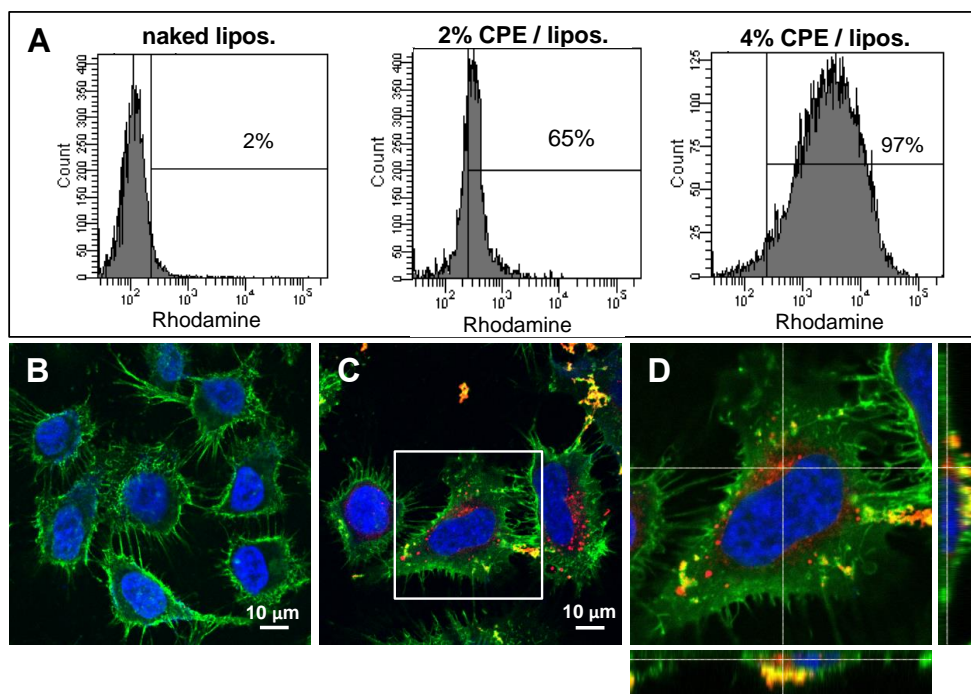


Figure 5. (A) Cytofluorimetric profile of HeLa cells incubated with naked liposomes (naked lipos.), liposomes decorated with 2 mol % (2% CPE/lipos.) and with 4 mol% (4% CPE/lipos.) of Arg₄-DAG in medium at pH 7.4. (B) Confocal microscopic images of HeLa cells incubated with fluorescently labelled naked liposomes and (C) liposomes decorated with 4 mol% of Arg₄-DAG in medium at pH 7.4. (D) Magnification of the white square in (C) and Z-stack projection. Cell nuclei were stained with DAPI (blue), membranes with WGA-AlexaFluor 633(green) and liposomes were fluorescently labelled with Rhodamine-DHPE (red).

The confocal microscopic analysis confirmed that the CPE decoration remarkably enhanced the liposome association to cells and showed the vesicles accessed to the HeLa cell cytoplasm (Figure 5 B-D). The cytoplasm disposition of liposomes was confirmed by the z-stack projection (Figure 5D) from image of Figure 5C, which highlighted that the liposomes (red spots) were

mainly located within the cytosol and disposed in proximity of the cell membrane (green) and not in the nucleus (blue).

3.5.2 pH-controlled association of CPE-decorated liposomes to cancer cells

CPE-decorated liposomes (4 mol% Arg₄-DAG/lipids) coated with the pH-sensitive co-polymer mPEG_{5kDa}-SDM₈ (1:1 Arg₄-DAG/mPEG_{5kDa}-SDM₈ molar ratio) were tested with HeLa cancer cells at pH 7.4 and 6.5, which mimic blood and extracellular tumour environments, respectively. HeLa cells were selected since they are a well-established cancer cell model. Anderson and co-workers extensively characterized the tumor generated with these cells. He showed that interstitial space of this tumor has pH 6.5 ± 0.2 .⁶⁰ Thus, we identified pH 6.5 a representative condition to investigate association of the pH responsive liposomes with this model cell line. Figure 6A and B shows that at pH 6.5 the cells displayed a 6-fold higher mean fluorescence intensity (MFI) with respect to pH 7.4. Almost all cells (99%) were fluorescently positive to liposomes at pH 6.5 while cells were mostly spare from interaction with pH responsive liposomes when incubated at pH 7.4. Studies showed that mPEG_{5kDa}-SDM₈-free CPE-decorated liposomes massively associated with the cells regardless the incubation pH (Figure 5A, and Figure S8 and S9 in ESI). Furthermore, control liposomes decorated with 4 mol% of commercial mPEG_{5kDa}-DSPE yielded negligible cell association at both pHs (Figure 6A).

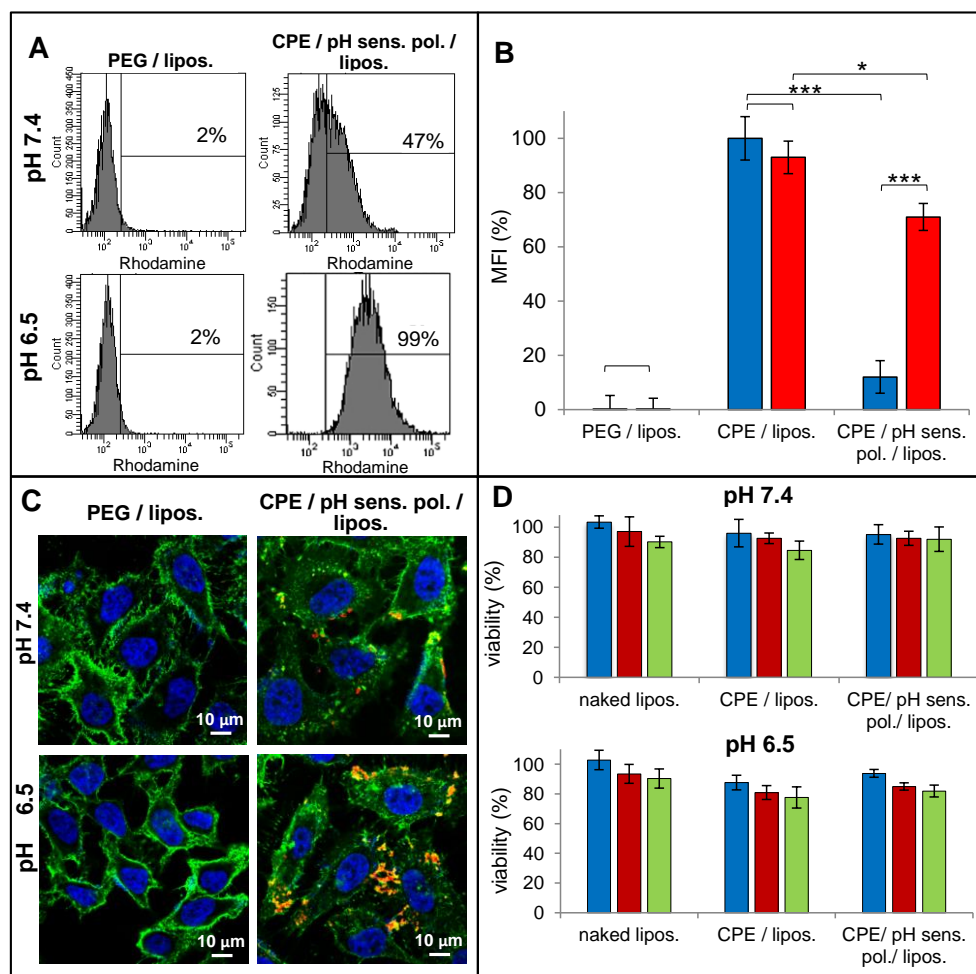


Figure 6. (A) Cytofluorimetric profile of HeLa cells incubated with rhodamine-DHPE labelled mPEG_{5kDa}-DSPE coated liposomes, and CPE-decorated liposomes with pH-responsive polymer in medium at pH 6.5 and 7.4. (B): relative MFI% of cells incubated with liposomes in medium at pH 7.4 (■) and 6.5 (■) derived from Figure (A). 100% relative MFI was attributed to cells incubated with CPE-decorated liposomes in medium at pH 7.4. Statistical analyses: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. (C) Confocal microscopic images of HeLa cells incubated with: mPEG_{5kDa}-DSPE coated liposomes, and CPE-decorated liposomes with pH-responsive polymer at pH 6.5 and 7.4. (D) Cytotoxicity profile of HeLa cells incubated with naked liposomes, CPE-

decorated liposomes with and without pH-responsive polymer in medium at pH 7.4 and 6.5 at 0.1 (■), 0.5 (■) and 1 (■) mg/mL for 6 hours.

The confocal microscopy images reported Figure 6C confirmed the pH controlled association of the mPEG_{5kDa}-SDM₈ coated Arg₄-DAG decorated liposomes with HeLa cells. The intense cell associated fluorescence detected at pH 6.5 was a consequence of the dissociation of mPEG_{5kDa}-SDM₈ from the Arg₄-DAG units on liposome surface which restored the CPE on the liposome surface yielding their preferential association to cells under acid conditions. On the contrary, at pH 7.4 only few red fluorescent spots were detected on the outer cell surface in proximity of cell membrane.

Oppositely to several cationic liposomes that display inherent high cytotoxicity⁶¹ Figure 6D shows that the Arg₄-DAG decorated liposomes possessed a negligible cell toxicity, which was similar to that of naked liposomes. Recent studies proved that cationic liposomes obtained by Arg decoration cause low cell toxicity when compared to the positive control Lipofectamine™ 2000. This result was attributed to the presence of amino acid analogs within the lipidic composition,⁶² which may provide a different interference of the liposomes with the membrane function and integrity of the cell or the subcellular compartments with respect to commercial lipids such as DOTAP.⁶³⁻⁶⁴ Notably, the presence of the pH-responsive polymer, both at pH 7.4 and 6.5, did not alter the biocompatibility of the CPE-decorated liposomes.

Interestingly, a control study carried out with human epithelial PNT2 cells, at pH 7.4, showed that under physiological conditions the pH responsive liposomes do not significantly interact with non-cancer cells (Figure S10). On the contrary, the CPE decorated liposomes showed an

unspecific remarkable interaction with the cells. This result further supports the environmental activation of the pH sensitive carrier, which can avoid interactions with normal cells under physiological conditions, namely at pH 7.4.

3.5.3 Enhancement and pH controlled delivery of liposome payload

Calcein and BSA loaded CPE-decorated liposomes were used to investigate the intracellular drug delivery.

Calcein was chosen as model hydrophilic molecule that does not cross lipid bilayers, thus it mimics drugs with negligible access to the intracellular compartment.⁶⁵ The calcein loaded CPE-decorated vesicles displayed a size of 187 ± 8 nm with a polydispersity index of 0.117, and the loading capacity and encapsulation efficiency were found to be 52 μ g calcein/mg lipids and 5.6%, respectively.

Proteins possess low permeability across biological membranes while the intracellular delivery can have therapeutic applications for those possessing cytosolic targets. Thus, a delivery system is required to promote the access of these macromolecules to the intracellular targets.⁶⁶ BSA was selected as model biomacromolecule since it was already exploited to study the delivery of proteins encapsulated in liposomes.⁶⁷ Proteins with surface hydrophobic regions, such as BSA,⁶⁸ may favorably associate with the lipidic components of liposomes during the loading process and thus be suitably encapsulated. Notably, a consensus exists concerning the favorable effect of the protein/lipid interaction on the protein loading efficiency in liposomes.⁶⁹ Although extensive BSA labelling was considered useful to emphasize the cell up-take results, in the present study,

BSA was labelled with a minimal amount of Rhodamine (3 Rhodamine molecules per BSA, Figure S11) in order to avoid structural protein alteration. BSA-Rho loaded CPE-decorated liposomes were obtained with a size of 201 ± 6 nm and a polydispersity index of 0.197. The slight size increase of BSA-Rho loaded liposomes with respect to their non-loaded counterparts was already observed in our previous work.⁶⁷ After removal of the non-loaded BSA-Rho, RP-HPLC analysis indicated a loading capacity of 25.2 μ g BSA-Rho/mg lipids and an encapsulation efficiency of 2.52 %.

The release study from CPE-decorated liposomes was carried out in buffer at pH 7.4 and pH 6.5, mimicking the blood and extracellular tumour environments, respectively (Figure 7).

Preliminary studies confirmed that calcein fluorescence is not affected by the pH. The study showed that the encapsulated calcein is not released by liposomes at the two pH conditions for at least 16 days, which is consistent with the low permeability of calcein across the lipid bilayer⁶⁵ and with the fact that the CPE does not alter the vesicle membrane permeability (Figure 7A).

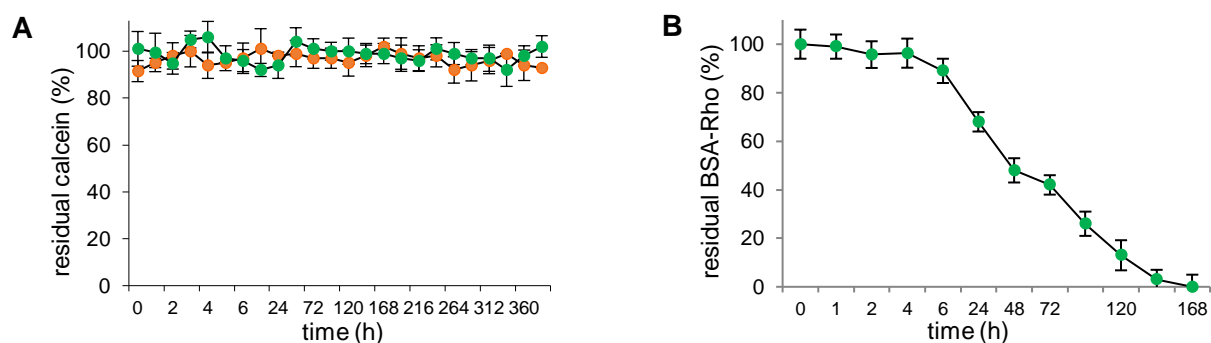


Figure 7. Calcein (**A**) and BSA-Rho (**B**) release profile from CPE-decorated liposomes in PBS at pH 7.4 (●) and 6.5 (●) at 25 °C.

The release profiles reported in Figure 7B indicate that the release of BSA-Rho is quite slow in the first 6 hours and that approximately 30% of the loaded protein was released in 24 h. Notably, BSA-Rho release was complete in about 7 days. The release profile was very similar to that observed for PEG decorated liposomes reported in our previous study.⁶⁷ To note that the decoration of liposomes with CPE, as well as the pH conditions (Figure S12), neither altered the loading capacity nor the release profile of the vesicles.

The enhancement of delivery of small hydrophilic molecules to cancer cells was tested with calcein loaded CPE-decorated liposomes. The results of the flow cytometric investigation reported in Figure 8A, showed the ability of the CPE-decorated vesicles to deliver calcein to HeLa cells while the free calcein and calcein loaded naked liposomes did not associate to the cells. All cells treated with CPE-decorated liposomes showed a higher fluorescence intensity with respect to controls at pH 7.4 and 6.5 (Figure 8 and S13 in ESI).

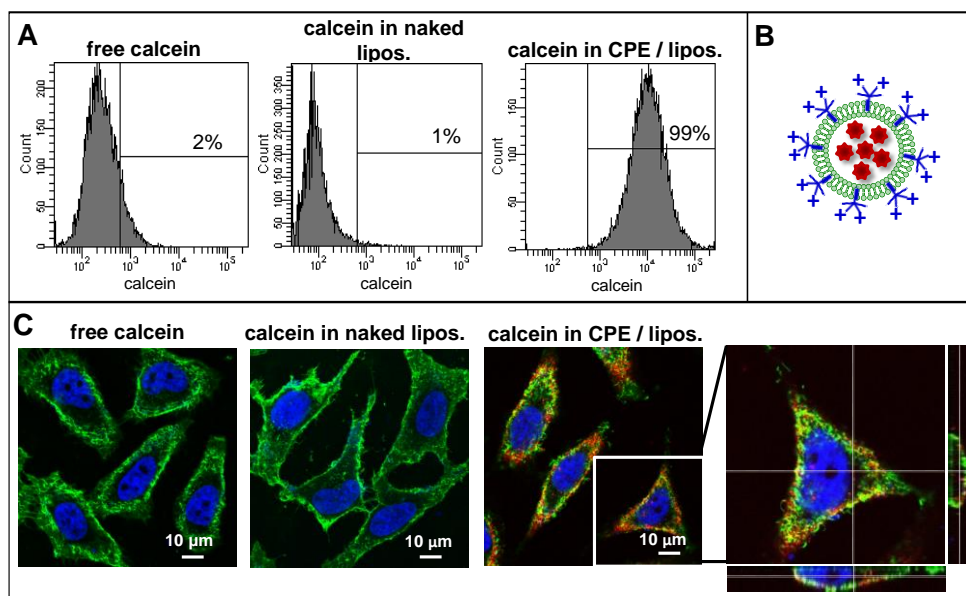


Figure 8. Cytofluorimetric profile (A) and confocal microscopic images (C) of HeLa cells incubated with free calcein, calcein loaded naked liposomes (calcein in naked lipos.) and

liposomes decorated with 4 mol% of Arg₄-DAG (calcein in CPE/lipos.) in medium at pH 7.4.

The last right image of panel **(C)** represents a magnification of the image of cells incubated with “calcein in CPE/lipos” and the corresponding z-stack. Cell nucleus was stained with DAPI in blue, cell membrane is represented in green upon staining with WGA-AlexaFluor 633, and calcein in red. Panel **(B)** schematically represents the calcein loaded liposomes.

Confocal microscopy confirmed the disposition of calcein within the cytosol when delivered by CPE-labelled liposomes (Figure 8C) while a very low cell associated fluorescence of calcein was detected neither when cells were incubated with equimolar concentration of free calcein, which was expected due to the limited cell membrane permeability of this molecule, nor when incubated with calcein loaded naked liposomes.

The cytofluorimetric data reported in Figure 9A and B show that 1 h incubation with BSA-Rho loaded CPE-decorated liposomes yielded 33% fluorescently positive cells and higher mean fluorescence intensity (MFI) compared to free BSA-Rho and BSA-Rho loaded naked liposomes that showed negligible cell association and MFI.

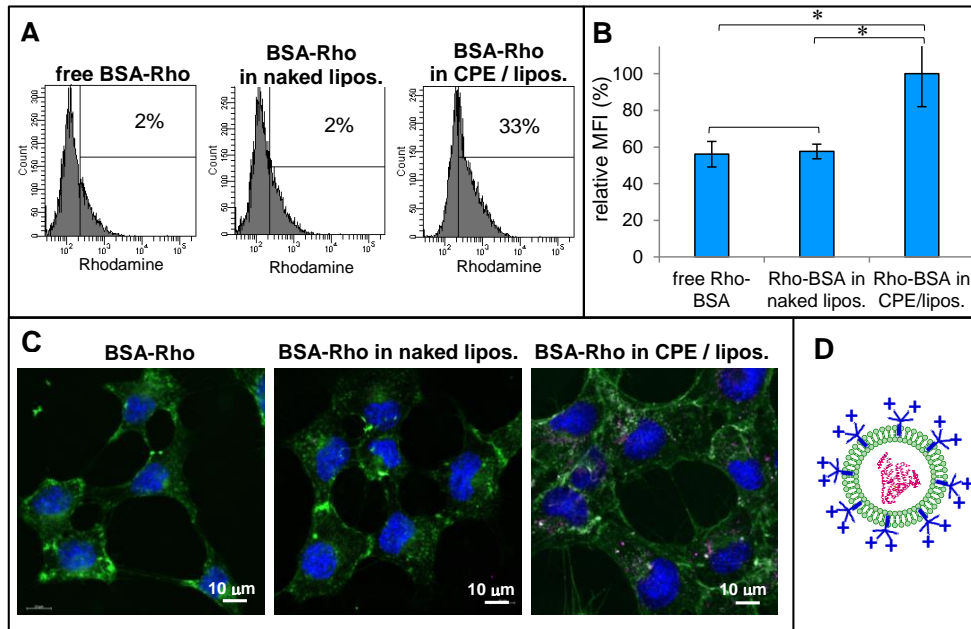


Figure 9. Cell association of BSA-Rho loaded liposomes with HeLa cells. **(A)** Cytofluorimetric profiles, **(B)** relative MFI%, and **(C)** confocal microscopic images of HeLa cells treated with free BSA-Rho, naked liposomes and CPE-decorated liposomes loaded with BSA-Rho in medium at pH 7.4. Panel **(D)** schematically represents CPE-decorated liposomes loaded with BSA-Rho. Statistical analyses: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

The confocal microscopy images reported in Figure 9C show that after cell incubation with BSA-Rho loaded CPE-decorated liposomes, BSA-Rho disposed intracellularly while no intracellular fluorescence was observed by cell incubation with free BSA-Rho and BSA-Rho loaded naked liposomes. These results prove that Arg₄-DAG decorated liposomes can efficiently provide for intracellular delivery of biomacromolecules.

The different cell up-take of BSA-Rhodamine and calcein loaded Arg₄-DAG decorated liposomes observed by cytofluorimetry and confocal microscopy (Figure 8 and 9) is ascribable to the different fluorescence quantum yield and content of the two fluorophores (rhodamine and calcein) in the formulations.

The pH controlled delivery of calcein to cancer cells was tested by incubating HeLa cells with calcein loaded CPE-decorated liposomes formulated with mPEG_{5kDa}-SDM₈ at pH 7.4 and 6.5. The cytofluorimetric analysis indicated that the pH-sensitive liposomes selectively delivered calcein to the cells under tumour acid environment. In fact, cells incubated with the pH responsive formulation at pH 7.4 (mimicking the blood) displayed a 6.8 times lower MFI with respect to cells incubated with the same formulation at pH 6.5 mimicking the tumour (Figure 10A and B).

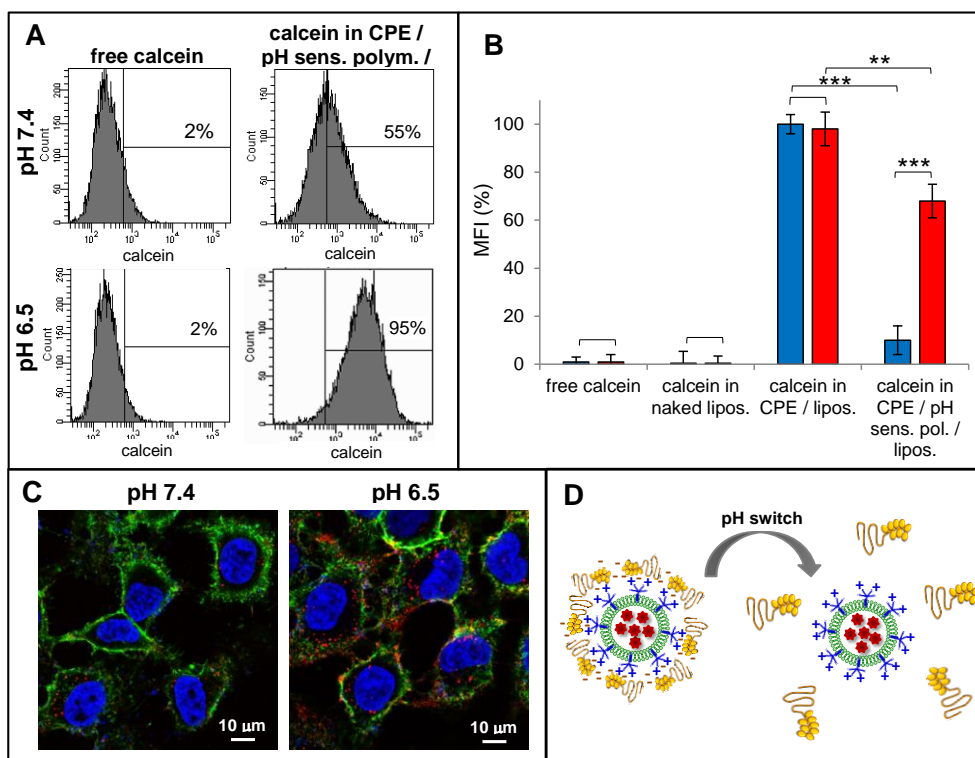


Figure 10. (A) Cytofluorimetric profiles of HeLa cells incubated with calcein-loaded CPE-decorated liposomes with and without mPEG_{5kDa}-SDM₈ in medium at pH 7.4 and 6.5 (Free

calcein was used as reference). **(B)** Relative MFI% of HeLa cells at pH 7.4 (■) and 6.5 (■) from panel **(A)**; statistical analyses: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. **(C)** Confocal microscopic images of HeLa cells treated with calcein loaded CPE-decorated liposomes with mPEG_{5kDa}-SDM₈ at pH 7.4 and 6.5 and schematic representation of the CPE unshielding process triggered at acid pH **(D)**.

The confocal imaging study (Figure 10C) also confirmed that, while calcein encapsulated in the CPE-decorated liposomes remarkably disposed in the cell cytosol regardless of the incubation pH (Figure 8 and S13 in ESI), the calcein intracellular delivery by the pH responsive liposomes was higher under acidic pH (6.5) with respect to pH 7.4 (Figure 10C).

These results confirmed that these pH-sensitive liposomes are efficient vehicles for controlled and preferential intracellular delivery of loaded molecules that, otherwise, would not access the cytosol across cell membranes and can be exploited for site-selective association to and uptake by cancer cells under the tumour matrix pH. Furthermore, the pH-guided intracellular delivery of drugs using these pH-sensitive liposomes may provide a valuable strategy to enhance the access to cancer cells after accumulation in the tumor while minimizing off-site toxicity to healthy tissues. Further studies are in progress to elucidate the intracellular trafficking of these liposomes and the payload release mechanism. Notably, evidences are reported in the literature that nanocarriers modified with CPPs appeared to enter the cytosol by endosomal escape.⁷⁰

4. Conclusions

We have investigated here a novel strategy for site-programmed delivery of therapeutic molecules to cancer cells by combining on liposome surface a synthetic cell penetration enhancer and a polymeric pH sensor. These components modulate the surface features of the liposome according to a dynamic behaviour. The pH responsive PEG shielding prevents the carrier from the interaction with cells under condition mimicking the physiological environment, namely blood and extracellular environment of healthy tissues, while revealing the cell penetration enhancer on the liposome surface which guides the access of the lipidic vesicles to the cytosol of cancer cells under environmental conditions that are significant for certain diseases such as those of the tumour matrix.

The responsiveness and efficacy of the system stems from the physicochemical features of the two poly-ionic components combined on the vesicle surface and from the strategy and ratio of assembly of these functional components. Therefore, the sequential decoration of liposomes with the CPE and the coating with the PEGylating masking agent has allowed the generation of “smart” vesicles for site-selective drug delivery.

The results reported in this work demonstrate that the environmentally controlled *shielding-and-revealing* strategy can provide for enhanced cancer cell delivery of drug loaded liposomes without using active biorecognition targeting. This strategy can be applied to a variety of nanocarriers to enhance site-selectivity to the tumour. Further studies will be undertaken to elucidate the pharmacokinetic and biodistribution behaviour of this system in vivo.

ASSOCIATED CONTENT

Supporting Information.

Supporting information include: chemicals and equipment; synthesis and characterization of the cell penetration enhancer (CPE), the pH-sensitive copolymer and the Rhodamine labelled BSA; methods to assess decoration efficiency of liposome with Arg₄-DAG and liposome cytotoxicity; methods for liposome immobilization on SPR sensorchips and additional SPR results; SAXS spectra of control liposomes; cell association of control liposomes by FACS and confocal microscopy.

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NOTES

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